



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/787,219
Source: JFW
Date Processed by STIC: 7/30/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

~~TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER~~
~~VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND~~
~~TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:~~

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box-1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
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IFWO

RAW SEQUENCE LISTING

DATE: 07/30/2004

PATENT APPLICATION: US/10/787,219

TIME: 11:40:38

Input Set : A:\248628US0X.txt

Output Set: N:\CRF4\07302004\J787219.raw

3 <110> APPLICANT: JESTIN, JEAN-LUC
 4 VICHIER-GUERRE, SOPHIE
 6 <120> TITLE OF INVENTION: METHODS FOR OBTAINING THERMOSTABLE ENZYMES, DNA POLYMERASE I
 7 VARIANTS FROM THERMUS AQUATICUS HAVING NEW CATALYTIC ACTIVITIES,
 8 METHODS FOR OBTAINING THE SAME, AND APPLICATIONS OF THE SAME
 10 <130> FILE REFERENCE: 248628USOX
 12 <140> CURRENT APPLICATION NUMBER: 10/787,219
 13 <141> CURRENT FILING DATE: 2004-02-27
 15 <160> NUMBER OF SEQ ID NOS: 61
 17 <170> SOFTWARE: PatentIn version 3.3
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 24
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Artificial Sequence
 24 <220> FEATURE:
 25 <223> OTHER INFORMATION: Synthetic DNA
 27 <400> SEQUENCE: 1
 28 taacaatagg ccggccaccc cttc
 31 <210> SEQ ID NO: 2
 32 <211> LENGTH: 18
 33 <212> TYPE: DNA
 34 <213> ORGANISM: Artificial Sequence
 36 <220> FEATURE:
 37 <223> OTHER INFORMATION: Synthetic DNA
 39 <400> SEQUENCE: 2
 40 gagtttttgt tctgcggc
 43 <210> SEQ ID NO: 3
 44 <211> LENGTH: 27
 45 <212> TYPE: DNA
 46 <213> ORGANISM: Artificial Sequence
 48 <220> FEATURE:
 49 <223> OTHER INFORMATION: Synthetic DNA
 51 <400> SEQUENCE: 3
 52 tttaatcatc tgcagtaccg ggagctc
 55 <210> SEQ ID NO: 4
 56 <211> LENGTH: 28
 57 <212> TYPE: DNA
 58 <213> ORGANISM: Artificial Sequence
 60 <220> FEATURE:
 61 <223> OTHER INFORMATION: Synthetic DNA
 63 <400> SEQUENCE: 4
 64 ttcattcttg ctagctcctg ggagaggc
 67 <210> SEQ ID NO: 5

Does Not Comply
Corrected Diskette Needed

pp 2, 10

24

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RAW SEQUENCE LISTING

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68 <211> LENGTH: 43
69 <212> TYPE: DNA
70 <213> ORGANISM: Artificial Sequence
72 <220> FEATURE:
73 <223> OTHER INFORMATION: Synthetic DNA
76 <220> FEATURE:
77 <221> NAME/KEY: misc_feature OK
78 <222> LOCATION: (15)..(15)
79 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
80 C and A in the trimer sequence CAR and AVY, respectively
82 <220> FEATURE:
83 <221> NAME/KEY: misc_feature
84 <222> LOCATION: (16)..(16)
85 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
86 A and V, in the trimer sequence CAR and AVY, respectively
88 <220> FEATURE:
89 <221> NAME/KEY: misc_feature
90 <222> LOCATION: (17)..(17)
91 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
92 R and Y, in the trimer sequence CAR and AVY, respectively
94 <220> FEATURE:
95 <221> NAME/KEY: misc_feature
96 <222> LOCATION: (24)..(24)
97 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
98 C and A, in the trimer sequence CAR and AVY, respectively
100 <220> FEATURE:
101 <221> NAME/KEY: misc_feature
102 <222> LOCATION: (25)..(25)
103 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
104 A and R, in the trimer sequence CAR and AVY, respectively
106 <220> FEATURE:
107 <221> NAME/KEY: misc_feature
108 <222> LOCATION: (26)..(26)
109 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
110 R and Y, in the trimer sequence CAR and AVY, respectively
112 <400> SEQUENCE: 5
W--> 113 cgggccaccc cttnnnctc aacnnncggg accagctgga aag 43
116 <210> SEQ ID NO: 6
117 <211> LENGTH: 65
118 <212> TYPE: DNA
119 <213> ORGANISM: Artificial Sequence
121 <220> FEATURE:
122 <223> OTHER INFORMATION: Synthetic DNA
125 <220> FEATURE:
126 <221> NAME/KEY: misc_feature
127 <222> LOCATION: (17)..(17)
128 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative

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abundance:

129 Y and R, in the trimer sequence ~~YTG~~ and ~~RRT~~, respectively
131 <220> FEATURE:

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132 <221> NAME/KEY: misc_feature

133 <222> LOCATION: (18)..(18)

134 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

135 T and B, in the trimer sequence YTG and RBT, respectively

137 <220> FEATURE:

138 <221> NAME/KEY: misc_feature

139 <222> LOCATION: (19)..(19)

140 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

141 G and T, in the trimer sequence YTG and RBT, respectively

143 <220> FEATURE:

144 <221> NAME/KEY: misc_feature

145 <222> LOCATION: (20)..(20)

146 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

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149 <220> FEATURE:

150 <221> NAME/KEY: misc_feature

151 <222> LOCATION: (21)..(21)

152 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

153 T and B, in the trimer sequence YTG and RBT, respectively

155 <220> FEATURE:

156 <221> NAME/KEY: misc_feature

157 <222> LOCATION: (22)..(22)

158 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

159 G and T, in the trimer sequence YTG and RBT, respectively

161 <220> FEATURE:

162 <221> NAME/KEY: misc_feature

163 <222> LOCATION: (26)..(26)

164 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

165 Y and R, in the trimer sequence YTG and RBT, respectively

167 <220> FEATURE:

168 <221> NAME/KEY: misc_feature

169 <222> LOCATION: (27)..(27)

170 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

171 T and B, in the trimer sequence YTG and RBT, respectively

173 <220> FEATURE:

174 <221> NAME/KEY: misc_feature

175 <222> LOCATION: (28)..(28)

176 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

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179 <220> FEATURE:

180 <221> NAME/KEY: misc_feature

181 <222> LOCATION: (44)..(44)

182 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

183 Y and R, in the trimer sequence YTG and RBT, respectively

185 <220> FEATURE:

186 <221> NAME/KEY: misc_feature

187 <222> LOCATION: (45)..(45)

188 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

189 T and B, in the trimer sequence YTG and RBT, respectively

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191 <220> FEATURE:
192 <221> NAME/KEY: misc_feature
193 <222> LOCATION: (46)..(46)
194 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
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197 <400> SEQUENCE: 6
W--> 198 ggatgaggtc cggcaannnn nnaatnnngg tgctcttcag cttnnngagc tcccgtact      60
200 gcagg                                         65
203 <210> SEQ ID NO: 7
204 <211> LENGTH: 62
205 <212> TYPE: DNA
206 <213> ORGANISM: Artificial Sequence
208 <220> FEATURE:
209 <223> OTHER INFORMATION: Synthetic DNA
212 <220> FEATURE:
213 <221> NAME/KEY: misc_feature
214 <222> LOCATION: (17)..(17)
215 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
216      C and A, in the trimer sequence CAR and AVY, respectively
218 <220> FEATURE:
219 <221> NAME/KEY: misc_feature
220 <222> LOCATION: (18)..(18)
221 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
222      A and V, in the trimer sequence CAR and AVY, respectively
224 <220> FEATURE:
225 <221> NAME/KEY: misc_feature
226 <222> LOCATION: (19)..(19)
227 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
228      R and Y, in the trimer sequence CAR and AVY, respectively
230 <220> FEATURE:
231 <221> NAME/KEY: misc_feature
232 <222> LOCATION: (32)..(32)
233 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
234      C and A, in the trimer sequence CAR and AVY, respectively
236 <220> FEATURE:
237 <221> NAME/KEY: misc_feature
238 <222> LOCATION: (33)..(33)
239 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
240      A and V, in the trimer sequence CAR and AVY, respectively
242 <220> FEATURE:
243 <221> NAME/KEY: misc_feature
244 <222> LOCATION: (34)..(34)
245 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative
abundance:
246      R and Y, in the trimer sequence CAR and AVY, respectively
248 <220> FEATURE:
249 <221> NAME/KEY: misc_feature

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250 <222> LOCATION: (41)..(41)

251 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

252 C and A, in the trimer sequence CAR and AVY, respectively

RAW SEQUENCE LISTING

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Input Set : A:\248628US0X.txt

Output Set : N:\CRF4\07302004\J787219.raw

254 <220> FEATURE:

255 <221> NAME/KEY: misc_feature

256 <222> LOCATION: (42)..(42)

257 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

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260 <220> FEATURE:

261 <221> NAME/KEY: misc_feature

262 <222> LOCATION: (43)..(43)

263 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

264 R and Y, in the trimer sequence CAR and AVY, respectively

266 <400> SEQUENCE: 7

W--> 267 caaccagacg gccacgnna cgggcaggct annagctcc nnncccaacc tccagaacat 60

269 cc 62

272 <210> SEQ ID NO: 8

273 <211> LENGTH: 43

274 <212> TYPE: DNA

275 <213> ORGANISM: Artificial Sequence

277 <220> FEATURE:

278 <223> OTHER INFORMATION: Synthetic DNA

281 <220> FEATURE:

282 <221> NAME/KEY: misc_feature

283 <222> LOCATION: (14)..(14)

284 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

285 Y and R, in the trimer sequence YTG and RBT, respectively

287 <220> FEATURE:

288 <221> NAME/KEY: misc_feature

289 <222> LOCATION: (15)..(15)

290 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

291 T and B, in the trimer sequence YTG and RBT, respectively

293 <220> FEATURE:

294 <221> NAME/KEY: misc_feature

295 <222> LOCATION: (16)..(16)

296 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

297 G and T, in the trimer sequence YTG and RBT, respectively

299 <220> FEATURE:

300 <221> NAME/KEY: misc_feature

301 <222> LOCATION: (23)..(23)

302 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

303 Y and R, in the trimer sequence YTG and RBT, respectively

305 <220> FEATURE:

306 <221> NAME/KEY: misc_feature

307 <222> LOCATION: (24)..(24)

308 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

309 T and B, in the trimer sequence YTG and RBT, respectively

311 <220> FEATURE:

312 <221> NAME/KEY: misc_feature

313 <222> LOCATION: (25)..(25)

314 <223> OTHER INFORMATION: n represents the following sequences in a 1:1 relative abundance:

315 G and T, in the trimer sequence YTG and RBT, respectively

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/787,219

DATE: 07/30/2004
TIME: 11:40:39

Input Set : A:\248628US0X.txt
Output Set: N:\CRF4\07302004\J787219.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 15, 16, 17, 24, 25, 26
Seq#:6; N Pos. 17, 18, 19, 20, 21, 22, 26, 27, 28, 44, 45, 46
Seq#:7; N Pos. 17, 18, 19, 32, 33, 34, 41, 42, 43
Seq#:8; N Pos. 14, 15, 16, 23, 24, 25
Seq#:9; N Pos. 20, 21, 22, 38, 39, 40, 44, 45, 46, 47, 48, 49
Seq#:10; N Pos. 20, 21, 22, 29, 30, 31, 44, 45, 46
Seq#:11; N Pos. 19, 20, 21, 28, 29, 30

VERIFICATION SUMMARY

DATE: 07/30/2004

PATENT APPLICATION: US/10/787,219

TIME: 11:40:39

Input Set : A:\248628US0X.txt

Output Set: N:\CRF4\07302004\J787219.raw

L:113 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:318 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:472 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:528 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0